

# DEN\_WT\_BR2

%features

60  
40  
20  
0

■ % in genome  
▨ detected  
■ % in sample

2.9  
1.9  
1  
0

protein\_coding  
pseudogene  
lincRNA  
antisense  
miRNA  
snoRNA  
snRNA  
IG\_V\_gene  
processed\_transcript  
misc\_RNA  
rRNA  
sense\_intronic  
IG\_J\_gene  
IG\_C\_gene  
non\_coding  
polymorphic\_pseudogene  
sense\_overlapping  
IG\_D\_gene  
3prime\_overlapping\_ncrna  
ncrna\_host

